1642



RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/451,739E

DATE: 08/07/2002 TIME: 13:07:32

```
Input Set : A:\EP.txt
                     Output Set: N:\CRF4\08072002\I451739E.raw
      1 <110> APPLICANT: Jager, Dirk
              Scanlan, Matthew
      2
      3
              Gure, Ali
      4
              Jager, Elke
              Knuth, Alexander
              Old, Lloyd
      6
              Chen, Yao-tseng
      7
      9 <120> TITLE OF INVENTION: Isolated Nucleic Acid Molecules Encoding Cancer Associated
Antigens,
              the Antigens per se, and Uses Thereof
     10
     12 <130> FILE REFERENCE: LUD 5615
     14 <140> CURRENT APPLICATION NUMBER: 09/451,739E
     16 <141> CURRENT FILING DATE: 1999-11-30
                                                                  ENTERED
     18 <160> NUMBER OF SEQ ID NOS: 19
     21 <210> SEQ ID NO: 1
     22 <211> LENGTH: 1533
     23 <212> TYPE: DNA
     24 <213> ORGANISM: Homo sapiens
W--> 25 <220> FEATURE:
     26 <221> NAME/KEY: CDS
     27 <222> LOCATION: 235
     28/<223> OTHER INFORMATION: unknown
     29 <400> SEQUENCE: 1
     30 ggttttecae gttggaeaag tgeggetegg eggeeagegg agegegeece tteeegetge
                                                                            120
     32 cogeteeget eetetettet acceageeea gtgggegagt gggeagegge ggeegeggeg
     34 ctgggccctc tcccgccggt gtgtgcgcgc tcgtacgcgc ggcccccggc gccagccccg
W--> 36 ccgcctgaga gggggcctgc gccgccggcc ggggcgtgcg cccggggagcc accgncaccg
     38 eggeeegege eeteaggege tggggteeee geggaeeegg aggeggegga egggetegge
                                                                            360
     40 agatqtagec geegggeega ageaggagee ggeggggggg egeegggaga gegagggett
     42 tgcattttgc agtgctattt tttgaggggg gcggagggtg gaggaagtcg gaaagccgcg
                                                                            420
     44 ccgagtcgcc ggggacctcc ggggtgaacc atgttgagtc ctgccaacgg ggagcagctc
                                                                            480
     46 cacctggtga actatgtgga ggactacctg gactccatcg agtccctgcc tttcgacttg
                                                                            540
                                                                            600
     48 cagagaaatg tetegetgat gegggagate gaegegaaat accaagagat eetgaaggag
     50 ctagacgagt gctacgagcg cttcagtcgc gagacagacg gggcgcagaa gcggcggatg
                                                                            660
     52 ctgcactgtg tgcagcgcgc gctgatccgc agccaggagc tgggcgacga gaagatccag
                                                                            720
                                                                            780
     54 atcgtgagcc agatggtgga gctggtggag aaccgcacgc ggcaggtgga cagccacgtg
     56 gagetgtteg aggegeagea ggagetggge gacacagegg geaacagegg caaggetgge
                                                                            840
     58 geggacagge ceaaaggega ggeggeageg caggetgaca ageceaacag caagegetea
     60 cggcggcage gcaacaacga gaaccgtgag aacgcgtcca gcaaccacga ccacgacgac
```

62 ggcgcctcgg gcacacccaa ggagaagaag gccaagacct ccaagaagaa gaagcgctcc 1020 64 aaggccaagg cggagcgaga ggcgtcccct gccgacctcc ccatcgaccc caacgaaccc 1080 66 acgtactgtc tgtgcaacca ggtctcctat ggggagatga tcggctgcga caacgacgag 1140 69 tgccccatcg agtggttcca cttctcgtgc gtggggctca atcataaacc caagggcaag 1200 71 tggtactgtc ccaagtgccg gggggagaac gagaagacca tggacaaagc cctggagaaa 1260

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```
73 tocaaaaaag agagggetta caacaggtag tttgtggaca ggcgcctggt gtgaggagga 1320
     75 caaaataaac cgtgtattta ttacattgct gcctttgttg aggtgcaagg agtgtaaaat 1380
     77 qtatattttt aaaqaatqtt agaaaaggaa ccattccttt catagggatg gcagtgattc 1440
     79 tqtttqcctt ttqttttcat tqqtacacqt qtaacaaqaa aqtqqtctqt qqatcaqcat 1500
     81 tttagaaact acaaatatag gtttgattca aca
     84 <210> SEQ ID NO: 2
     85 <211> LENGTH: 1143
     86 <212> TYPE: DNA
     87 <213> ORGANISM: Homo sapiens
W--> 88 <400> SEQUENCE: 2
     89 gagtaacceg ataatatgee gttgteegge aeggegaega gaatteeeag atatageagt
     91 agcagtgate eegggeetgt ggetegggge eggggetgea gtteggaeeg eeteeegega
                                                                            120
     93 cccgcggggg ctcggagaca gtttcaggcc gcatctttgc tgacccgagg gtggggccgc
                                                                            180
     95 gcgtggccgt ggaaacagat cctgaaggag ctagacgagt gctacgagcg cttcagtcgc
                                                                            240
     97 gagacagacg gggcgcagaa gcggcggatg ctgcactgtg tgcagcgcgc gctgatccgc
     99 agccaggagc tgggcgacga gaagatccag atcgtgagcc agatggtgga gctggtggag
     101 aaccgcacgc ggcaggtgga cagccacgtg gagctgttcg aggcgcagca ggagctgggc
                                                                             420
                                                                             480
    103 gacacagtgg gcaacagcgg caaggttggc gcggacaggc ccaatggcga tgcggtagcg
                                                                             540
     105 caqtctqaca aqcccaacaq caaqcqctca cqqcqqcaqc qcaacaacqa qaaccqtqaq
    107 aacgcqtcca qcaaccacqa ccacqacqac qqcqcctcqq qcacacccaa qqaqaaqaaq
                                                                             600
    109 gccaagacct ccaagaagaa gaagcgctcc aaggccaagg cggagcgaga ggcgtcccct
                                                                             660
    111 geogaectee ceategaece caacgaacce acgtactgte tgtgcaacca ggteteetat
                                                                            720
    113 qqqqaqatqa tcqqctqcqa caacqacqaq tgccccatcq agtggttcca cttctcgtgc
                                                                            780
    115 qtqqqqctca atcataaacc caaqqqcaag tggtactgtc ccaagtgccg gggggagaac
                                                                            840
    117 gagaagacca tggacaaagc cctggagaaa tccaaaaaag agagggctta caacaggtag
    119 tttgtggaca ggcgcctggt gtgaggagga caaaataaac cgtgtattta ttacattgct
    121 gcctttgttg aggtgcaagg agtgtaaaat gtatatttt aaagaatgtt agaaaaggaa 1020
    123 ccattccttt catagggatg gcagtgattc tgtttgcctt ttgttttcat tggtacacgt 1080
    125 gtaacaagaa agtggtctgt ggatcagcat tttagaaact acaaatatag gtttgattca 1140
                                                                            1143
    127 aca
    130 <210> SEQ ID NO: 3
    131 <211> LENGTH: 742
    132 <212> TYPE: DNA
     133 <213> ORGANISM: Homo sapiens
W--> 134 <220> FEATURE:
W--> 135 <400> SEQUENCE: 3
    137 cgccgtccac accccagcgg ccctgacgct gtcccctccg cgaccctcgc ctctggaaaa
                                                                              60
                                                                             120
    139 aqtqacaqqc aaqqccacqc cccqcqaqq gccqqcctcq agcccqcaqc ccccagggcc
    141 tgggacgaga tcctgaagga gctagacgag tgctacgagc gcttcagtcg cgagacagac
                                                                             240
    143 gqqqcqcaqa aqcqqcqqat qctqcactqt qtqcaqcqcq cqctqatccq cagccaggag
                                                                             300
    145 ctgggcgacg agaagatcca gatcgtgagc cagatggtgg agctggtgga gaaccgcacg
    147 cggcaggtgg acagccacgt ggagctgttc gaggcgcagc aggagctggg cgacacagcg
                                                                             360
                                                                             420
    149 ggcaacagcg gcaaggctgg cgcggacagg cccaaaggcg aggcggcagc gcaggctgac
                                                                             480
    151 aagcccaaca gcaagcgctc acggcggcag cgcaacaacg agaaccgtga gaacgcgtcc
                                                                             540
    153 agcaaccacg accacgacga cggcgcctcg ggcacaccca aggagaagaa ggccaagacc
    155 tecaagaaga agaagegete caaggeeaag geggagegag aggegteeee tgeegaeete - 600
    157 cccatcgacc ccaacgaacc cacgtactgt ctgtgcaacc aggtctccta tggggagatg
                                                                             660
                                                                             720
    159 atcggctgcg acaacgacga gtgccccatc gagtggttcc acttctcgtg cgtggggctc
    161 aatcataaac ccaagggcaa gt
                                                                             742
```

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Input Set : A:\EP.txt

Output Set: N:\CRF4\08072002\I451739E.raw

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164 <210> SEQ ID NO: 4
165 <211> LENGTH: 857
166 <212> TYPE: DNA
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167 <213> ORGANISM: Homo sapiens

W--> 168 <400> SEQUENCE: 4

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169 cctccgagaa cggtgtccat ggcacagggc gggaagagat aaggcctagg gaaggcgccc
171 ctcgggccta tccacctctt ctggggctcg gcactaggaa gcagcttccc tctcaggccc
                                                                        120
173 ctttgtctcc aagccgttcc aaactgagta ccgggagacg acacaaaggg agggcggtga
                                                                       180
175 cggatggcgc aggcgcggga gccgcctagg ctgctgggag tggtggtccg gccgcggaat
                                                                       240
177 ggagatectg aaggagetag aegagtgeta egagegette agtegegaga eagaegggge
                                                                       300
179 gcaqaagegg eggatgetge actgtgtgea gegegegetg ateegeagee aggagetggg
                                                                       360
181 cgacgagaag atccagatcg tgagccagat ggtggagctg gtggagaacc gcacgcggca
                                                                       420
183 ggtggacagc cacgtggagc tgttcgaggc gcagcaggag ctgggcgaca cagcgggcaa
                                                                        480
185 cageggeaag getggegegg acaggeceaa aggegaggeg geagegeagg etgacaagee
                                                                        540
187 caacagcaag cgctcacggc ggcagcgcaa caacgagaac cgtgagaacg cgtccagcaa
                                                                       600
                                                                       660
189 ccacgaccac gacgacggcg cctcgggcac acccaaggag aagaaggcca agacctccaa
191 gaagaagaag cgctccaagg ccaaggcgga gcgagaggcg tcccctgccg acctccccat
                                                                       720
193 cgaccccaac gaacccacgt actgtctgtg caaccaggtc tcctatgggg agatgatcgg
                                                                       780
195 ctgcgacaac gacgagtgcc ccatcgagtg gttccacttc tcgtgcgtgg ggctcaatca
                                                                       840
197 taaacccaaq qqcaaqt
                                                                        857
```

205 <210> SEQ ID NO: 5 206 <211> LENGTH: 279 207 <212> TYPE: PRT

208 <213> ORGANISM: Homo sapiens

W--> 209 <400> SEQUENCE: 5

210 Met Leu Ser Pro Ala Asn Gly Glu Gln Leu His Leu Val Asn Tyr Val 211 1 213 Glu Asp Tyr Leu Asp Ser Ile Glu Ser Leu Pro Phe Asp Leu Gln Arg 214 20 25 216 Asn Val Ser Leu Met Arg Glu Ile Asp Ala Lys Tyr Gln Glu Ile Leu 219 Lys Glu Leu Asp Glu Cys Tyr Glu Arg Phe Ser Arg Glu Thr Asp Gly 55 222 Ala Gln Lys Arg Arg Met Leu His Cys Val Gln Arg Ala Leu Ile Arg 70 225 Ser Gln Glu Leu Gly Asp Glu Lys Ile Gln Ile Val Ser Gln Met Val 85 90 228 Glu Leu Val Glu Asn Arg Thr Arg Gln Val Asp Ser His Val Glu Leu 105 100 231 Phe Glu Ala Gln Glu Leu Gly Asp Thr Val Gly Asn Ser Gly Lys 232 120 234 Val Gly Ala Asp Arg Pro Asn Gly Asp Ala Val Ala Gln Ser Asp Lys 135 140 237 Pro Asn Ser Lys Arg Ser Arg Gln Arg Asn Asn Glu Asn Arg Glu

240 Asn Ala Ser Ser Asn His Asp His Asp Asp Gly Ala Ser Gly Thr Pro

243 Lys Glu Lys Lys Ala Lys Thr Ser Lys Lys Lys Arg Ser Lys Ala

155

170

150

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244

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Input Set : A:\EP.txt

Output Set: N:\CRF4\08072002\I451739E.raw

```
246 Lys Ala Glu Arg Glu Ala Ser Pro Ala Asp Leu Pro Ile Asp Pro Asn
                195
                                     200
     249 Glu Pro Thr Tyr Cys Leu Cys Asn Gln Val Ser Tyr Gly Glu Met Ile
                                215
     252 Gly Cys Asp Asn Asp Glu Cys Pro Ile Glu Trp Phe His Phe Ser Cys
                            230
                                                235
     255 Val Gly Leu Asn His Lys Pro Lys Gly Lys Trp Tyr Cys Pro Lys Cys
                        245
                                            250
     258 Arg Gly Glu Asn Glu Lys Thr Met Asp Lys Ala Leu Glu Lys Ser Lys
                    260
                                         265
     261 Lys Glu Arg Ala Tyr Asn Arg
                275
     262
     265 <210> SEQ ID NO: 6
     266 <211> LENGTH: 210
     267 <212> TYPE: PRT
     268 <213> ORGANISM: Homo sapiens
W--> 269 <220> FEATURE:
W--> 270 <400> SEQUENCE: 6
     273 Met Leu His Cys Val Gln Arg Ala Leu Ile Arg Ser Gln Glu Leu Gly
     276 Asp Glu Lys Ile Gln Ile Val Ser Gln Met Val Glu Leu Val Glu Asn
                                        25
                    20
     279 Arg Thr Arg Gln Val Asp Ser His Val Glu Leu Phe Glu Ala Gln Gln
                                    40
     282 Glu Leu Gly Asp Thr Val Gly Asn Ser Gly Lys Val Gly Ala Asp Arg
                                55
     285 Pro Asn Gly Asp Ala Val Ala Gln Ser Asp Lys Pro Asn Ser Lys Arg
     288 Ser Arg Arg Gln Arg Asn Asn Glu Asn Arg Glu Asn Ala Ser Ser Asn
                        85
                                            90
     291 His Asp His Asp Asp Gly Ala Ser Gly Thr Pro Lys Glu Lys Lys Ala
                                        105
     294 Lys Thr Ser Lys Lys Lys Arg Ser Lys Ala Lys Ala Glu Arg Glu
                                    120
     297 Ala Ser Pro Ala Asp Leu Pro Ile Asp Pro Asn Glu Pro Thr Tyr Cys
                                135
         130
     300 Leu Cys Asn Gln Val Ser Tyr Gly Glu Met Ile Gly Cys Asp Asn Asp
                                                155
    303 Glu Cys Pro Ile Glu Trp Phe His Phe Ser Cys Val Gly Leu Asn His
                        165
     306 Lys Pro Lys Gly Lys Trp Tyr Cys Pro Lys Cys Arg Gly Glu Asn Glu
     307
                    180
                                        185
     309 Lys Thr Met Asp Lys Ala Leu Glu Lys Ser Lys Lys Glu Arg Ala Tyr
                                    200
     312 Asn Arg
    313
         210
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316 <210> SEQ ID NO: 7 317 <211> LENGTH: 235 318 <212> TYPE: PRT RAW SEQUENCE LISTING DATE: 08/07/2002 PATENT APPLICATION: US/09/451,739E TIME: 13:07:32

Input Set : A:\EP.txt

Output Set: N:\CRF4\08072002\I451739E.raw

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319 <213> ORGANISM: Homo sapiens
W--> 320 <400> SEQUENCE: 7
     321 Met Glu Ile Leu Lys Glu Leu Asp Glu Cys Tyr Glu Arg Phe Ser Arg
     324 Glu Thr Asp Gly Ala Gln Lys Arg Arg Met Leu His Cys Val Gln Arg
                     20
                                         25
     327 Ala Leu Ile Arg Ser Gln Glu Leu Gly Asp Glu Lys Ile Gln Ile Val
                 35
                                     40
     330 Ser Gln Met Val Glu Leu Val Glu Asn Arg Thr Arg Gln Val Asp Ser
                                 55
     333 His Val Glu Leu Phe Glu Ala Gln Gln Glu Leu Gly Asp Thr Val Gly
                             70
                                                 75
     336 Asn Ser Gly Lys Val Gly Ala Asp Arg Pro Asn Gly Asp Ala Val Ala
     337
     341 Gln Ser Asp Lys Pro Asn Ser Lys Arg Ser Arg Arg Gln Arg Asn Asn
     342
                     100
                                         105
     344 Glu Asn Arg Glu Asn Ala Ser Ser Asn His Asp His Asp Asp Gly Ala
                 115
                                     120
     347 Ser Gly Thr Pro Lys Glu Lys Lys Ala Lys Thr Ser Lys Lys Lys
                                 135
                                                     140
     350 Arg Ser Lys Ala Lys Ala Glu Arg Glu Ala Ser Pro Ala Asp Leu Pro
                             150
                                                 155
     353 Ile Asp Pro Asn Glu Pro Thr Tyr Cys Leu Cys Asn Gln Val Ser Tyr
                         165
                                             170
     356 Gly Glu Met Ile Gly Cys Asp Asn Asp Glu Cys Pro Ile Glu Trp Phe
                                         185
     357
     359 His Phe Ser Cys Val Gly Leu Asn His Lys Pro Lys Gly Lys Trp Tyr
                 195
     362 Cys Pro Lys Cys Arg Gly Glu Asn Glu Lys Thr Met Asp Lys Ala Leu
                                 215
     365 Glu Lys Ser Lys Lys Glu Arg Ala Tyr Asn Arg
     366 225
                             230
     369 <210> SEQ ID NO: 8
     370 <211> LENGTH: 772
     371 <212> TYPE: DNA
     372 <213> ORGANISM: Homo sapiens
W--> 373 <220> FEATURE:
     374 <221> NAME/KEY: CDS
     375 <222> LOCATION: 689,714
     376 <223> OTHER INFORMATION: Unknown
W--> 377 <400> SEQUENCE: 8
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     380 cccacgacga cgtcacctcg ggcacgccca aggagaagaa agcccagacc tctaagaaga
                                                                            120
     382 agcagggete catggecaag gegtagegge aggegteece egcagacete eccategace
                                                                            180
     384 ccagcgagcc ctcctactgg gagatgatcc gctgcgacaa cgaatgcccc atcgagtggt
     386 tecgettete gtgtgtgagt etcaaccata aaccaaageg caagtggtae tgtteeagat 300
     388 gccggggaaa gaacgatggg caaagccctt gagaagtcca gaaaaaaaac agggcttata 360
     390 acaggtagtt tggggacatg cgtctaatag tgaggagaac aaaataagcc agtgtgttga 420
     392 ttacattgcc acctttgctg aggtgcagga agtgtaaaat gtatattttt aaagaatgtt 480
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/451,739E

DATE: 08/07/2002 TIME: 13:07:33

Input Set : A:\EP.txt

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 235 Seq#:8; N Pos. 689,714

Seq#:15; N Pos. 1628,1752,1758,1769,1789,1873,1908,1915,1933,1970,1976,2022